ATGGCGTCACAGAAGAGACCCTCCCAGAGGCACGGATCCAAGTACCTGGCCACAGCAAGTACCATGGACC
1
M A S Q K R P S Q R H G S K Y L A T A S T M D

ATGCCAGGCATGGCTTCCTCCCAAGGCACAGAGACACGGGCATCCTTGACTCCATCGGGCGCTTCTTTGG

H A R H G F L P R H R D T G I L D S I G R F F G

CGGTGACAGGGGTGCGCCAAAGCGGGGCTCTGGCAAGGACTCACACCACCCGGCAAGAACTGCTCACTAT

G D R G A P K R G S G K D S H H P A R T A H Y

GGCTCCCTGCCCCAGAAGTCACACGGCCGGACCCAAGATGAAAACCCCGTAGTCCACTTCTTCAAGAACA

G S L P Q K S H G R T Q D E N P V V H F F K N

TTGTGACGCCTCGCACACCACCCCCGTCGCAGGGAAAGGGGAGAGGACTGTCCCTGAGCAGATTTAGCTG

I V T P R T P P P S Q G K G R G L S L S R F S W

GGGGGCCGAAGGCCAGAGACCAGGATTTGGCTACGGAGGCAGAGCGTCCGACTATAAATCGGCTCACAAG

G A E G Q R P G F G Y G G R A S D Y K S A H K

GGATTCAAGGGAGTCGATGCCCAGGGCACGCTTTCCAAAATTTTTAAGCTGGGAGGAAGAGATAGTCGCT

G F K G V D A Q G T L S K I F K L G G R D S R

CTGCATCAGCCATCGCTAGACGCTGA   (SEQ ID NO: 4)

Fig.